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Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: Mon Sep 24 11:16:20 EDT 2007

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Application No: 10580567

Version No: 1.0

Input Set:

Output Set:

Started: 2007-09-12 15:59:18.228

Finished: 2007-09-12 15:59:18.926

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 698 ms

Total Warnings: 8

Total Errors: 0

No. of SeqIDs Defined: 10

Actual SeqID Count: 10

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SEQUENCE LISTING

<110> NAKAMURA, Koji et al.

<120> METHOD OF DETECTING LIVER CANCER, DIAGNOSTIC FOR LIVER
CANCER AND REMEDY FOR CANCER

<130> 0760-0355PUS1

<140> 10580567

<141> 2007-09-12

<150> PCT/JP2004/017499

<151> 2004-11-25

<160> 10

<170> PatentIn version 3.1

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agccggaccc gcgcccgcac cgctcccggg accgcgaccc cggccgcca gag atg 176
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acc gcg acc gaa gcc ctc ctg cgc gtc ctc ttg ctc ctg ctg gct ttc 224
Thr Ala Thr Glu Ala Leu Leu Arg Val Leu Leu Leu Leu Ala Phe
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ggc cac agc acc tat ggg gct gaa tgc ttc ccg gcc tgc aac ccc caa 272
Gly His Ser Thr Tyr Gly Ala Glu Cys Phe Pro Ala Cys Asn Pro Gln
20 25 30

aat gga ttc tgc gag gat gac aat gtt tgc agg tgc cag cct ggc tgg 320
Asn Gly Phe Cys Glu Asp Asp Asn Val Cys Arg Cys Gln Pro Gly Trp
35 40 45

cag ggt ccc ctt tgt gac cag tgc gtg acc tct ccc ggc tgc ctt cac 368
Gln Gly Pro Leu Cys Asp Gln Cys Val Thr Ser Pro Gly Cys Leu His
50 55 60 65

gga ctc tgt gga gaa ccc ggg cag tgc att tgc acc gac ggc tgg gac 416
Gly Leu Cys Gly Glu Pro Gly Gln Cys Ile Cys Thr Asp Gly Trp Asp
70 75 80

ggg gag ctc tgt gat aga gat gtt cgg gcc tgc tcc tcg gcc ccc tgt	464
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gcc aac aac ggg acc tgc gtg agc ctg gac ggt ggc ctc tat gaa tgc	512
Ala Asn Asn Gly Thr Cys Val Ser Leu Asp Gly Gly Leu Tyr Glu Cys	
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tcc tgt gcc ccc ggg tac tcg gga aag gac tgc cag aaa aag gac ggg	560
Ser Cys Ala Pro Gly Tyr Ser Gly Lys Asp Cys Gln Lys Lys Asp Gly	
115 120 125	
ccc tgt gtg atc aac ggc tcc ccc tgc cag cac gga ggc acc tgc gtg	608
Pro Cys Val Ile Asn Gly Ser Pro Cys Gln His Gly Gly Thr Cys Val	
130 135 140 145	
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Asp Asp Glu Gly Arg Ala Ser His Ala Ser Cys Leu Cys Pro Pro Gly	
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Phe Ser Gly Asn Phe Cys Glu Ile Val Ala Asn Ser Cys Thr Pro Asn	
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cca tgc gag aac gac ggc gtc tgc act gac att ggg ggc gac ttc cgc	752
Pro Cys Glu Asn Asp Gly Val Cys Thr Asp Ile Gly Gly Asp Phe Arg	
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Cys Arg Cys Pro Ala Gly Phe Ile Asp Lys Thr Cys Ser Arg Pro Val	
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Thr Asn Cys Ala Ser Ser Pro Cys Gln Asn Gly Gly Thr Cys Leu Gln	
210 215 220 225	
cac acc cag gtg agc tac gag tgt ctg tgc aag ccc gag ttc aca ggt	896
His Thr Gln Val Ser Tyr Glu Cys Leu Cys Lys Pro Glu Phe Thr Gly	
230 235 240	
ctc acc tgt gtc aag aag cgc gcg ctg agc ccc cag cag gtc acc cgt	944
Leu Thr Cys Val Lys Lys Arg Ala Leu Ser Pro Gln Gln Val Thr Arg	
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Leu Pro Ser Gly Tyr Gly Leu Ala Tyr Arg Leu Thr Pro Gly Val His	
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Glu Leu Pro Val Gln Gln Pro Glu His Arg Ile Leu Lys Val Ser Met	
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Lys Glu Leu Asn Lys Lys Thr Pro Leu Leu Thr Glu Gly Gln Ala Ile	
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Cys Phe Thr Ile Leu Gly Val Leu Thr Ser Leu Val Val Leu Gly Thr	
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Val Gly Ile Val Phe Leu Asn Lys Cys Glu Thr Trp Val Ser Asn Leu	
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Arg Tyr Asn His Met Leu Arg Lys Lys Asn Leu Leu Leu Gln Tyr Asn	
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Ser Gly Glu Asp Leu Ala Val Asn Ile Ile Phe Pro Glu Lys Ile Asp	
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Trp Gln Gly Pro Leu Cys Asp Gln Cys Val Thr Ser Pro Gly Cys Leu	
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Asp Gly Glu Leu Cys Asp Arg Asp Val Arg Ala Cys Ser Ser Ala Pro	

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Cys Ala Asn Asn Gly Thr Cys Val Ser Leu Asp Gly Gly Leu Tyr Glu
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Cys Ser Cys Ala Pro Gly Tyr Ser Gly Lys Asp Cys Gln Lys Lys Asp
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Val Thr Asn Cys Ala Ser Ser Pro Cys Gln Asn Gly Gly Thr Cys Leu
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Gly Leu Thr Cys Val Lys Lys Arg Ala Leu Ser Pro Gln Gln Val Thr
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Arg Leu Pro Ser Gly Tyr Gly Leu Ala Tyr Arg Leu Thr Pro Gly Val
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His Glu Leu Pro Val Gln Gln Pro Glu His Arg Ile Leu Lys Val Ser
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Met Lys Glu Leu Asn Lys Lys Thr Pro Leu Leu Thr Glu Gly Gln Ala
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Thr Val Gly Ile Val Phe Leu Asn Lys Cys Glu Thr Trp Val Ser Asn
325 330 335

Leu Arg Tyr Asn His Met Leu Arg Lys Lys Asn Leu Leu Leu Gln Tyr
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